



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/618,126
Source: ORP
Date Processed by STIC: 7/28/2003

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PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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1. EFS-Bio (<http://www.uspto.gov/efc/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 04/24/2003



OIPF

RAW SEQUENCE LISTING

DATE: 07/28/2003

PATENT APPLICATION: US/10/618,126

TIME: 13:36:19

Input Set : A:\MSB-7295.ST25.txt

Output Set: N:\CRF4\07282003\J618126.raw

3 <110> APPLICANT: Bayer Pharmaceuticals Corporation
 4 FROLAND, Wayne
 5 KELNER, Drew
 6 DUMAS, Michael
 7 PAN, Clark
 8 WHELAN, James
 9 WANG, John
 10 WANG, Wei
 12 <120> TITLE OF INVENTION: PITUITARY ADENYLATE CYCLASE ACTIVATING PEPTIDE (PACAP)
 RECEPTOR 3
 13 (VPAC2) AGONISTS AND THEIR PHARMACOLOGICAL METHODS OF USE
 15 <130> FILE REFERENCE: MSB-7295
 17 <140> CURRENT APPLICATION NUMBER: US/10/618,126
 17 <141> CURRENT FILING DATE: 2003-07-11
 17 <150> PRIOR APPLICATION NUMBER: US 60/395,738
 18 <151> PRIOR FILING DATE: 2002-07-12
 20 <160> NUMBER OF SEQ ID NOS: 264
 22 <170> SOFTWARE: PatentIn version 3.2

ERRORED SEQUENCES

39 <210> SEQ ID NO: 2
 40 <211> LENGTH: 31
 41 <212> TYPE: PRT
 42 <213> ORGANISM: Homo sapiens
 45 <220> FEATURE:
 46 <221> NAME/KEY: MISC_FEATURE
 47 <222> LOCATION: (1)..(31)
 48 <223> OTHER INFORMATION: Ac is acetyl
 50 <400> SEQUENCE: 2
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 E--> 53 1 5 10 15
 56 Val Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr
 57 20 25 30
 615 <210> SEQ ID NO: 40
 616 <211> LENGTH: 31
 617 <212> TYPE: PRT
 618 <213> ORGANISM: Homo sapiens
 621 <220> FEATURE:
 622 <221> NAME/KEY: MISC_FEATURE
 623 <222> LOCATION: (1)..(31)
 624 <223> OTHER INFORMATION: Ac is acetyl
 626 <400> SEQUENCE: 40

Does Not Comply
 Corrected Diskette Needed

pp 1-7

delete this - do not show these
 in the sequence.

Please explain
 modification
 in 22207-22237
 section without
 including "Ac"

p.2

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Input Set : A:\MSB-7295.ST25.txt

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E--> 628 Ac-His Thr Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
E--> 629      1          5          10          15
632 Val Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr
633      20          25          30
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1192 <211> LENGTH: 31
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1198 <221> NAME/KEY: MISC_FEATURE
1199 <222> LOCATION: (1)..(31)
1200 <223> OTHER INFORMATION: Ac is acetyl
1202 <400> SEQUENCE: 78
E--> 1204 Ac-His Thr Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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1208 Val Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Asn Lys Arg Tyr
1209      20          25          30
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1753 <211> LENGTH: 32
1754 <212> TYPE: PRT
1755 <213> ORGANISM: Homo sapiens
1758 <220> FEATURE:
1759 <221> NAME/KEY: MISC_FEATURE
1760 <222> LOCATION: (1)..(32)
1761 <223> OTHER INFORMATION: PEG is polyethylene glycol
1763 <400> SEQUENCE: 115
1765 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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1770      20          25          30
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1775 <212> TYPE: PRT
1776 <213> ORGANISM: Homo sapiens
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1780 <221> NAME/KEY: MISC_FEATURE
1781 <222> LOCATION: (1)..(32)
1782 <223> OTHER INFORMATION: Ac is acetyl; PEG is polyethylene glycol
1784 <400> SEQUENCE: 116
E--> 1786 Ac-His Thr Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
E--> 1787      1          5          10          15
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1791      20          25          30
1794 <210> SEQ ID NO: 117
1795 <211> LENGTH: 32
1796 <212> TYPE: PRT
1797 <213> ORGANISM: Homo sapiens
1800 <220> FEATURE:
1801 <221> NAME/KEY: MISC_FEATURE

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*same
type of
error*

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Input Set : A:\MSB-7295.ST25.txt

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1802 <222> LOCATION: (1)..(32)
1803 <223> OTHER INFORMATION: PEG is polyethylene glycol
1805 <400> SEQUENCE: 117
1807 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1808 1 5 10 15
E--> 1811 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr Cys-PEG
1812 20 25 30
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1816 <211> LENGTH: 30
1817 <212> TYPE: PRT
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1822 <221> NAME/KEY: MISC_FEATURE
1823 <222> LOCATION: (1)..(30)
1824 <223> OTHER INFORMATION: PEG is polyethylene glycol
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1828 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1829 1 5 10 15
E--> 1832 Val Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Cys-PEG
1833 20 25 30
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1837 <211> LENGTH: 32
1838 <212> TYPE: PRT
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1843 <221> NAME/KEY: MISC_FEATURE
1844 <222> LOCATION: (1)..(32)
1845 <223> OTHER INFORMATION: PEG is polyethylene glycol
1847 <400> SEQUENCE: 119
1849 His Thr Glu Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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E--> 1853 Val Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr Cys-PEG
1854 20 25 30
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1858 <211> LENGTH: 32
1859 <212> TYPE: PRT
1860 <213> ORGANISM: Homo sapiens
1863 <220> FEATURE:
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1865 <222> LOCATION: (1)..(32)
1866 <223> OTHER INFORMATION: PEG is polyethylene glycol
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1870 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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E--> 1874 Leu Ala Val Lys Lys Tyr Leu Gln Asp Ile Lys Gln Gly Gly Thr Cys-PEG
1875 20 25 30
1878 <210> SEQ ID NO: 121
1879 <211> LENGTH: 31
1880 <212> TYPE: PRT

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RAW SEQUENCE LISTING

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Input Set : A:\MSB-7295.ST25.txt

Output Set: N:\CRF4\07282003\J618126.raw

1881 <213> ORGANISM: Homo sapiens
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1885 <221> NAME/KEY: MISC_FEATURE
1886 <222> LOCATION: (1)..(31)
1887 <223> OTHER INFORMATION: PEG is polyethylene glycol
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1891 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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E--> 1895 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Cys-PEG
1896 20 25 30
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1901 <212> TYPE: PRT
1902 <213> ORGANISM: Homo sapiens
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1907 <222> LOCATION: (1)..(32)
1908 <223> OTHER INFORMATION: PEG is polyethylene glycol
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1912 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1913 1 5 10 15
E--> 1916 Leu Ala Ala Lys Lys Tyr Leu Gln Thr Ile Lys Gln Lys Arg Tyr Cys-PEG
1917 20 25 30
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1921 <211> LENGTH: 32
1922 <212> TYPE: PRT
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1926 <220> FEATURE:
1927 <221> NAME/KEY: MISC_FEATURE
1928 <222> LOCATION: (1)..(32)
1929 <223> OTHER INFORMATION: PEG is polyethylene glycol
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1933 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1934 1 5 10 15
E--> 1937 Met Ala Ala Lys Lys Tyr Leu Gln Thr Ile Lys Gln Lys Arg Tyr Cys-PEG
1938 20 25 30
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1942 <211> LENGTH: 32
1943 <212> TYPE: PRT
1944 <213> ORGANISM: Homo sapiens
1947 <220> FEATURE:
1948 <221> NAME/KEY: MISC_FEATURE
1949 <222> LOCATION: (1)..(32)
1950 <223> OTHER INFORMATION: PEG is polyethylene glycol
1952 <400> SEQUENCE: 124
1954 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1955 1 5 10 15
E--> 1958 Met Ala Ala His Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr Cys-PEG
1959 20 25 30

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Input Set : A:\MSB-7295.ST25.txt

Output Set: N:\CRF4\07282003\J618126.raw

1962 <210> SEQ ID NO: 125
1963 <211> LENGTH: 32
1964 <212> TYPE: PRT
1965 <213> ORGANISM: Homo sapiens
1968 <220> FEATURE:
1969 <221> NAME/KEY: MISC_FEATURE
1970 <222> LOCATION: (1)..(32)
1971 <223> OTHER INFORMATION: PEG is polyethylene glycol
1973 <400> SEQUENCE: 125
1975 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1976 1 5 10 15
E--> 1979 Met Ala Ala Lys His Tyr Leu Gln Ser Ile Lys Gln Lys Arg Tyr Cys-PEG
1980 20 25 30
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1984 <211> LENGTH: 31
1985 <212> TYPE: PRT
1986 <213> ORGANISM: Homo sapiens
1989 <220> FEATURE:
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1991 <222> LOCATION: (1)..(31)
1992 <223> OTHER INFORMATION: PEG is polyethylene glycol
1994 <400> SEQUENCE: 126
1996 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
1997 1 5 10 15
E--> 2000 Met Ala Gly Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Cys-PEG
2001 20 25 30
2004 <210> SEQ ID NO: 127
2005 <211> LENGTH: 31
2006 <212> TYPE: PRT
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2010 <220> FEATURE:
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2012 <222> LOCATION: (1)..(31)
2013 <223> OTHER INFORMATION: PEG is polyethylene glycol
2015 <400> SEQUENCE: 127
2017 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2018 1 5 10 15
E--> 2021 Met Ala Lys Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Cys-PEG
2022 20 25 30
2025 <210> SEQ ID NO: 128
2026 <211> LENGTH: 31
2027 <212> TYPE: PRT
2028 <213> ORGANISM: Homo sapiens
2031 <220> FEATURE:
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2033 <222> LOCATION: (1)..(31)
2034 <223> OTHER INFORMATION: PEG is polyethylene glycol
2036 <400> SEQUENCE: 128
2038 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln

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RAW SEQUENCE LISTING

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Input Set : A:\MSB-7295.ST25.txt

Output Set: N:\CRF4\07282003\J618126.raw

2039 1 5 10 15
E--> 2042 Met Ala Arg Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Cys-PEG
2043 20 25 30
2046 <210> SEQ ID NO: 129
2047 <211> LENGTH: 31
2048 <212> TYPE: PRT
2049 <213> ORGANISM: Homo sapiens
2052 <220> FEATURE:
2053 <221> NAME/KEY: MISC_FEATURE
2054 <222> LOCATION: (1)..(31)
2055 <223> OTHER INFORMATION: PEG is polyethylene glycol
2057 <400> SEQUENCE: 129
2059 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2060 1 5 10 15
E--> 2063 Met Ala Ser Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Arg Cys-PEG
2064 20 25 30
2067 <210> SEQ ID NO: 130
2068 <211> LENGTH: 31
2069 <212> TYPE: PRT
2070 <213> ORGANISM: Homo sapiens
2073 <220> FEATURE:
2074 <221> NAME/KEY: MISC_FEATURE
2075 <222> LOCATION: (1)..(31)
2076 <223> OTHER INFORMATION: PEG is polyethylene glycol
2078 <400> SEQUENCE: 130
2080 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2081 1 5 10 15
E--> 2084 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Pro Gln Lys Arg Cys-PEG
2085 20 25 30
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2091 <213> ORGANISM: Homo sapiens
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2095 <221> NAME/KEY: MISC_FEATURE
2096 <222> LOCATION: (1)..(31)
2097 <223> OTHER INFORMATION: PEG is polyethylene glycol
2099 <400> SEQUENCE: 131
2101 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2102 1 5 10 15
E--> 2105 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Gln Gln Lys Arg Cys-PEG
2106 20 25 30
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2110 <211> LENGTH: 31
2111 <212> TYPE: PRT
2112 <213> ORGANISM: Homo sapiens
2115 <220> FEATURE:
2116 <221> NAME/KEY: MISC_FEATURE
2117 <222> LOCATION: (1)..(31)

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2118 <223> OTHER INFORMATION: PEG is polyethylene glycol
2120 <400> SEQUENCE: 132
2122 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2123 1 5 10 15
E--> 2126 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Arg Gln Lys Arg Cys-PEG
2127 20 25 30
2130 <210> SEQ ID NO: 133
2131 <211> LENGTH: 31
2132 <212> TYPE: PRT
2133 <213> ORGANISM: Homo sapiens
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2137 <221> NAME/KEY: MISC_FEATURE
2138 <222> LOCATION: (1)..(31)
2139 <223> OTHER INFORMATION: PEG is polyethylene glycol
2141 <400> SEQUENCE: 133
2143 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
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E--> 2147 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Arg Arg Cys-PEG
2148 20 25 30
2151 <210> SEQ ID NO: 134
2152 <211> LENGTH: 31
2153 <212> TYPE: PRT
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2159 <222> LOCATION: (1)..(31)
2160 <223> OTHER INFORMATION: PEG is polyethylene glycol
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2164 His Ser Asp Ala Val Phe Thr Asp Gln Tyr Thr Arg Leu Arg Lys Gln
2165 1 5 10 15
E--> 2168 Met Ala Ala Lys Lys Tyr Leu Gln Ser Ile Lys Gln Lys Ala Cys-PEG
2169 20 25 30

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/618,126

DATE: 07/28/2003

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Input Set : A:\MSB-7295.ST25.txt

Output Set: N:\CRF4\07282003\J618126.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No
 L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:52 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:52 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
 L:53 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2
 L:628 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:628 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
 L:629 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:40
 L:1204 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1204 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
 L:1205 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:78
 L:1769 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1786 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1786 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
 L:1787 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:116
 L:1790 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1811 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1832 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1853 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1874 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1895 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1916 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1937 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1958 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:1979 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2000 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2021 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2042 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2063 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2084 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2105 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2126 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2147 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2168 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2189 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2210 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2231 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2252 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2273 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2294 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2315 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2336 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2357 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2378 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2399 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2420 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2441 M:333 E: Wrong sequence grouping, Amino acids not in groups!
 L:2462 M:333 E: Wrong sequence grouping, Amino acids not in groups!

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Input Set : A:\MSB-7295.ST25.txt

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L:2483 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:2504 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:2525 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:2546 M:333 E: Wrong sequence grouping, Amino acids not in groups!